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Athersys, Inc.
3201 Carnegie Ave, Cleveland, (
Tel: 216 431 9900
Fax: 216 361 9596
Email: scain@athersys.com
High quality sequence stop: 49
Location/Qualifiers
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AW913564 Uf54e02.y
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AV226010 AV226010
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/Organism="Homo sapiens"
/Ob_xref="taxon:9606"
/Clone_lib="Athersys RAGE Library"
/Cell_line="HT1080"
/note="See 'Creation of Genome-wide Protein Expression
/Intaries using Random Activation of Gene Expression',
Nature Biotechnology, in press. Note that even though the
cell type indicated is HT1080, since a random activation
method was used, these sequence tags are not necessarily
expressed in HT1080 under normal circumstances."
211 a 195 c 178 g 222 t 1 others
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on,J.J., Sherf,B.,
Leventhal,C., Thor
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Tel:
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216 431 9900
216 361 9596
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quality sequence sto
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193 c 173 g 214 t 1 others
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/db_xref-"taxon:9606"
/clone_lib-"Athersys RAGE Library"
/cell_line-"HT1080"
/note-"See 'Creation of Genome-wide Protein Expression
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Pred. No. 4.6e-150;
0; Mismatches 3;
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ATCATCACCTATCTTTACTGGCAGATCACAGAGGGAAGGATGATGATGATAAGGCTGCTC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 This sequence was derived from the FAPESP/LICR Human Cancer Genome Project. This entry can be seen in the following URL (http://www.ludwig.org.br/scripts/gethtml2.pl?t1=QV4&t2=QV4-ST0212-120100-075-f09&t3=2000-01-12&t4=1)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Unpublished (1999)
Contact: Simpson A.J.G.
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EST.
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Ludwig Institute for Cancer Research
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The FAPESP/LICE Human Cancer Genome Project
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae;
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+55-11-2707001
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                                                                                                                                                                                                                                                                                                                                                                                                                                                       /note="Organ: stomach; Vector: puc18; Site_1: SmaI; Site_2: SmaI; A mini-library was made by cloning products derived from ORESTES PCR (U.S. Letters Patent application No. 196,716 - Ludwig Institute for Cancer Research) profiles into the pUC 18 vector. Reverse transcription of tissue mRNA and cDNA amplification were performed under Low stringency conditions."
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        /organism="Homo sapiens"
/db_xref="taxon:9606"
/clone_lib="ST0212"
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Location/Qualifiers
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Fax: 216 361 9596
                                                                                                                                                                                                                                                                                                                                                                                                                                                             Athersys, Inc.
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                                                                                                                                                                                                                                                                                                /clone_lib="Athersys RAGE Library"
/cell_line="HT1080"
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/db_xref="taxon:9606"
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98.2%;
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                                                                                                                                   Athersys, Inc.
3201 Carnegie Ave, (
Tel: 216 431 9900
Fax: 216 361 9596
                                                                                                                                                                                                                                                                                             Homo sapiens
Eukaryota; Me
Mammalia; Eut
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RST41079 Atl
BG221275
BG221275.1
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Creation of genome-wide protein expression libraries using random activation of gene expression
activation of gene expression
                                                                                                                                                                                                                                                                                                                                       EST
                                                                                                                                                                               Contact: Scott J. Cain
                                                                                                                                                                                                                                                                                                                             human
                                                                                                                                                                                                                                                                                mmalia; Eutheria; (bases 1 to 789)
                                                                                                     l: scain@athersys.com
quality sequence stop: :
Location/Qualifiers
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/organism="Homo sapiens"
/db_xref="taxon:9606"
/db_xref="taxon:9606"
/clone_lib="Athersys RAGE Library"
/cell_line="HT1080"
/note="See 'Creation of Genome-wide Protein Expression / Ibraries using Random Activation of Gene Expression', Libraries using Random Activation of Gene Expression', Nature Biotechnology, in press. Note that even though the cell type indicated is HT1080, since a random activation method was used, these sequence tags are not necessarily
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RAGE Library I
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RESULT 6
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Best Local Similarity 97.0%;
Matches 553; Conservative
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               Eukaryota; mermates; current Mammalla; Eutheria; Primates; current Mammalla; Eutheria; Primates; current 1 (bases 1 to 837)

RS Harrington,J.J., Sherf,B., Rundlett,S., Jackson,P.D., Perry,R., RS, Lerner,L., Costanzo,D., McElligott,R., Boozer,S., Mays,R., Smith, Lerner,L., Costanzo,D., McElligott,R., Boozer,S., Mays,R., Smith, E., Veloso,N., Klika,A., Hess,J., Cothren,K., Lo,K., Offenbacher,J., Danzig,J. and Ducar,M.

E Creation of genome-wide protein expression libraries using random activation of gene expression

B Creation of gene expression

B Creation of gene expression
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Pred. No. 2.1e-141;
0; Mismatches 16;
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Carnegie

Cleveland,

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44115,

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RESULT 7
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1 209 c 184 g 235 t 1 others
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/cell_line="HT1080"
/note="See 'Creation of Genome-wide Protein Expression
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/db_xref="taxon:9606"
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No. 7.2e-140;
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3201 Carnegie Ave, (
3201 Carnegie Ave, (
Tel: 216 431 9900
Fax: 216 319596
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Creation of genome-wide protein expression libraries using random activation of gene expression
activation of gene expression
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/note="See 'Creation of Genome-wide Protein Expression
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: 216 431 9900
: 216 361 9596
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/db_xref-"taxon:9606"
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/cell_line-"HT1080"
/note-"See 'Creation of Genome-wide Protein Expression
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Pred. No. 4.7e-138;
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                                                                                                                                                                                                                                                                                                                                                                  Email: asimpson@ludwig.org.br
Email: asimpson@ludwig.org.br
This sequence was derived from the FAPESP/LICR Human Cancer Genome
This sequence was derived from the following URL
This sequence was derived from the following URL
This sequence was derived from the FAPESP/LICR Human Cancer Genome
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QV4-ST0212-261199-045-c09
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261199-045-c09&t3=1999-11-26&t4=1)
Seq_primer: puc_18_forward
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Ludwig Institute for Cancer Research
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                   ρ
                                                                      /note="Organ: stomach; Vector: puc18; Site_1: SmaI; Site_2: SmaI; A mini-library was made by cloning products derived from ORESTES PCR (U.S. Letters Patent application No. 196,716 - Ludwig Institute for Cancer Research) profiles into the pUC 18 vector. Reverse transcription of
                   tissue mRNA and cDNA amplification were performed low stringency conditions." 173 c 164 g 184 t
                                                                                                                                                                    /clone_lib="ST0212"
/dev_stage="Adult"
                                                                                                                                                                                                                                                              Location/Qualifiers
                                                                                                                                                                                                         /db_xref="taxon:9606"
                                                                                                                                                                                                                               /organism="Homo sapiens"
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                                                                       Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalla; Eutheria; Primates; Catarrhini; Hominidae; Homo.

1 (bases 1 to 822)

Harrington,J.J., Sherf,B., Rundlett,S., Jackson,P.D., Perry,R., Cain,S., Leventhal,C., Thornton,M., Ramachandran,R., Whittington,J., Leventhal,C., Thornton,M., Ramachandran,R., Whittington,J., Lerner,L., Costanzo,D., McElligott,K., Boozer,S., Mays,R., Smith,E., Veloso,N., Kilka,A., Hess,J., Cothren,K., Lo,K., Offenbacher,J., Danzig,J. and Ducar,M.

Creation of genome-wide protein expression libraries using random activation of gene expression
Nat. Biotechnol. 19 (5), 440-445 (2001)
           Athersys, Inc.
3201 Carnegie Ave, Cleveland,
Tel: 216 431 9900
Fax: 216 361 9596
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RST39061 Athersys RAGE Library Homo sapiens cDNA,
BG219307
BG219307.1 GI:13745328
                                                      Contact: Scott J. Cain
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scain@athersys.com
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93.4%;
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Local Similarity 99.1 nes 446; Conservative
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RST18188 Ath
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BG198917.1
Creation of genome-wide protein expression libraries using random activation of gene expression Nat. Biotechnol. 19 (5), 440-445 (2001)
                                        Homo sapiens

Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;

Mammalia; Butheria; Primates; Catarrhini; Hominidae; Homo.

1 (bases 1 to 871)

1 (bases 1 to 871)

Harrington, J.J., Sherf, B., Rundlett, S., Jackson, P.D., Perry, R., Cain, S., Leventhal, C., Thornton, M., Ramachandran, R., Whittington, J., Lerner, L., Costanzo, D., McElligott, K., Boozer, S., Mays, R., Smith, E., Veloso, N., Kika, A., Hess, J., Cothren, K., Lo, K., Offenbacher, J., Danzig, J. and Ducar, M.

Creation of genome-wide protein expression libraries using random
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Location/Qualifiers
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/db_xref="taxon:9606"
/clone_llb="Athersys RAGE Library"
/cell_line="HT1080"
/note="see 'Creation of Genome-wide Protein Expression / note="see 'Creation of Genome-wide Protein Expression',
Libraries using Random Activation of Gene Expression',
Nature Biotechnology, in press. Note that even though the cell type indicated is HT1080, since a random activation method was used, these sequence tags are not necessarily expressed in HT1080 under normal circumstances."
41 a 169 c 179 g 230 t 3 others
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Contact: S
Athersys,
3201 Carne
Tel: 216 4
clone:4932443L08.
              clone_lib:RIKEN full-length
                             HTC; CAP trapper.
Mus musculus (str
                                                             AK016573.1 GI:12855380
                                                                                          PLACE1009493,
                                                                                                        Mus musculus adult male testis cDNA, RIKEN full-length enriched library, clone:4932443L08:homolog to CDNA FLJ13593 FIS, CLONE
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Libraries using Random Activation of Gene Expression', 
Nature Biotechnology, in press. Note that even though the 
cell type indicated is HT1080, since a random activation 
method was used, these sequence tags are not necessarily 
expressed in HT1080 under normal circumstances."
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/cell_line="HT1080"
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/db_xref="taxon:9606"
            (strain:C57BL/6J) adult male testis cDNA t
KEN full-length enriched mouse cDNA library
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Hayashizaki,...

Direct Submission

Research (RIKEN), Laboratory for Gen

Physical and Chemical Research (RIKEN), Laboratory for Gen

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Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
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ATTTTCTTGCTCTTCCCCATCCTTCACGGGAGTGCTGTGCACCCTGGCTATCACCATC
                   ATCTTCTTGCTCTTTTTCCCATCTTTCACCGGGGTCTTGTGCACCCTGGCCATCACCATC
                                                                            with BamHI and XhoI. cDNA of size comprised between 3 and 7 ki selected before cloning. Vector: a modified pBluescript KS(+) bulk excision from Lambda FLC I. Cloning sites, 5' end: SalI; and BamHI updt on 1100
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URL:http://genome.gsc.riken.go.jp/, Tel:81-45-503-9222,
Fax:81-45-503-9216)
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SSSAAGRVDCQIFNFGNDRTHALLRFSSLNESISQIYHGSECLTVDESCTFHETVQG
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SLSLTYRRTRNGLSELLNYITLWQKRFKVIGGKFGTSVLSYFSFLRWLLKFNITSFVM
MFSFILIPQFTVGAKNTLQFTGLEFFTGAGYFGDTVMYYGFTNSTIRHRMGGASYNM
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AEYNSEFLKTHRNPGAVLLLPFVVSCINLAVPRFYSMFRLVERYEIPRQEVYYLLVRN
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/clone_lib="RIKEN full-length enriched mouse cDNA library"
/dev_stage="adult"
207. ..2480
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LIGMKFTSLSLQEFDIARNVLELIYAQTLTWLGIFFCPLLPFIQMITLFIMFYVKNVS
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homolog to CDNA FLJ13593
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/strain="057BL/6J"
/db_xref="FANTOM_DB:4932443L08"
/db_xref="MGD:MGI:1907597"
/db_xref="taxon:10090"
/clone="4932443L08"
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/db_xref="GI:12855381"
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Creation of genome-wide protein expression libraries using random activation of gene expression.
                                                                                                                                                                             Email: scain@athersys.com
High quality sequence stop:
Location/Qualifiers
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3201 Carnegie Ave,
                                                                                                                                                                                                                                                                                                        Contact: Scott J. Cain
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/note="See 'Creation of Genome-wide Protein Expression Libraries using Random Activation of Gene Expression', Nature Biotechnology, in press. Note that even though the cell type indicated is HT1080, since a random activation method was used, these sequence tags are not necessarily
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                         453;
Email: asimpson@ludwig.org.br
This sequence was derived from t
Project. This entry can be seen
                                                           Laboratory of Cancer Genetics
Ludwig Institute for Cancer Research
Rua Prof. Antonio Prudente 109, 4 andar,
                                                                                                                                                     1 (bases 1 to 447)
Dias Neto, E., Garcia Correa, R., Verjovski-Almeida, S., Briones, M.R., Nagai, M.A., da Silva, W. Jr., Zago, M.A., Bordin, S., Costa, F.F., Goldman, G. H., Carvalho, A.F., Matsukuma, A., Baia, G.S., Simpson, D.H., Brunstein, A., deOliveira, P.S., Bucher, P., Jongeneel, C.V., O'Hare, M.J., Soares, F., Brentani, R.R., Reis, L.F., de Souza, S.J. and
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CM4-CN0089-130201-723-e04
BG982234
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Mammalia; I
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                                                                                            Contact: Simpson A.J.G.
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Pred. No. 3.5e-106;
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TCTCTTCATTCACTCCATCTACAGCTGGATCGACACCCTAAGTACACGGCCTGGCTACCT
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AW469181
AW469181.1
EST.
                                                                                                                                                                                                                                                                          AW469181
hc78f11.x1 I
similar to
                                      Tumor Gene Index Unpublished (1997)
Contact: Robert Strausberg,
Email: cgapbs-r@mail.nih.gov
                                                                             NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.
National Cancer Institute, Cancer Genome Anatomy
                                                                                                                    Eukaryota; Metazoa; Mammalia; Eutheria; 1 (bases 1 to 594)
                                                                                                                                                                               Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Seq primer: puc 18 forward
High quality sequence start: 25
High quality sequence stop: 447
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (http://www.ludwig.org.br/scripts/gethtml2.pl7t1=CM4&t2=CM4-CN0089-
130201-723-e04&t3=2001-02-13&t4=1)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               /note="Organ: colon_normal; Vector: puc18; Site_1: SmaI; Site_2: SmaI, A mini-library was made by cloning products derived from ORESTES PCR (U.S. Letters Patent application No. 196,716 - Ludwig Institute for Cancer Research) profiles into the pure lifector. Reverse transcription of tissue mRNA and cDNA pure lification were performed under low stringency conditions."
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            /organism="Homo sapiens"
/db_xref="taxon:9606"
/clone_lib="CN0089"
/dev_stage="Adult"
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1. .447
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                                                                                                                                                                                                                                        GI:7039287
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                                                                                                                                      Chordata;
Primates;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Score 401.4; DB 13
Pred. No. 2.4e-103;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Mismatches
                                                                                                                                                                                                                                                                                                                   594 bp
                                                                                                                                      Craniata; Vertebrata; Catarrhini; Hominidae
                                                                                                                                                                                                                                                                                                op mRNA linear EST 24-FEB-2000 sapiens cDNA clone IMAGE:2898861 3'
                                                                                                                                                                                                                                                                              :
                                                                                                                                                                                                                                                                                mRNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                         570
                                                                                                                                          Hominidae;
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                                                                                                                                                            Euteleostomi;
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Search completed: November Job time: 2224 secs
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              source
                                                                                                                                                                                      409 AAATTGATCAAGCTGCAGGATATGGAGAAGGAAAGCAAACCCCAGCTCACTTGTTCTGGAA 468
                                                                                                                     529 TTGCGATCTAGAAGATCAGTTCAAGAAGGTAATCCAAGGGCC 570
                                                                                                                                                                                                                                                                                                                                                                                    474 ATAAGGCTGCTCCATGAGCAGATCATTAATGAGGGCAAAGATAAAATGTTCCTGATAGAA 415
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                349 ATAAGGCTGCTCCATGAGCAGATCATTAATGAGGGCAAAGATAAAATGTTCCTGATAGAA 408
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      229 TGGGTTGTTTGGATCTATCGGAACCTCATTGGAAGTGTGCACTTCTTTTTCATCCTCACC 288
                                                                                              294 TTGCGATCTAGAAGATCAGTTCAAGAAGGTAATCCAAGGGCC 253
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               594 TGGGTTGTTGGAATCTATCGGAACCTCATTGGAAGTGTGCACTTCTTTTTCATCCTCACC 535
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CDNA Library Preparation: Life Technologies, Inc.
CDNA Library Arrayed by: Greg Lennon, Ph.D.
DNA Sequencing by: Washington University Genome Sequencing Center
Clone distribution: NCI-CGAP clone distribution information can be
found through the I.M.A.G.E. Consortium/LLNL at:
www-bio.llnl.gov/bbrp/image/image.html
Seq primer: -40Up from Gibco
High quality sequence stop: 404.
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Tissue Procurement: Christopher Moskaluk, M.D., Ph.D., Michael R. Emmert-Buck, M.D., Ph.D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    161 a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    /clone="IMAGE:2898861"
/clone_lib="NCI_CGAP_Gas4"
/tissue_type="poorly differentiated adenocarcinoma with
signet ring cell features"
/lab_host="DH10B"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        //note="Organ: stomach; Vector: pCMV-SPORT6; Site_1: SalI;
Site_2: NotI; Cloned unidirectionally. Primer: Oligo dT.
Average insert size 1.69 kb. Life Technologies catalog #:
11549-011"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            organism="Homo sapiens"
/db_xref="taxon:9606"
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99.4%;
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                        9, 2002, 03:15:54

 Mismatches

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